

# Use of *in vivo*-induced antigen technology to identify bacterial genes expressed during *Solea senegalensis* infection with *Photobacterium damsela* subsp. *piscicida*



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## Abstract

The marine fish pathogen *Photobacterium damsela* subsp. *piscicida* (*Phdp*) is responsible for important outbreaks affecting several fish species including flatfish *Solea senegalensis*. The aim of this work was to identify *in vivo*-induced expressed immunogenic proteins using pooled sera from fish that have experienced photobacteriosis. *In vivo* induced immunogenic proteins included inosine-5'-monophosphate dehydrogenase (Impdh) and alkyl hydroperoxide reductase (AhpC), two proteins involved in peptide synthesis: serine hydroxymethyl transferase (Shmt) and alanyl-tRNA synthetase (AlaRS) and the non-ribosomal peptide synthetase involved in the synthesis of the siderophore piscibactin (Irp2).

## Introduction

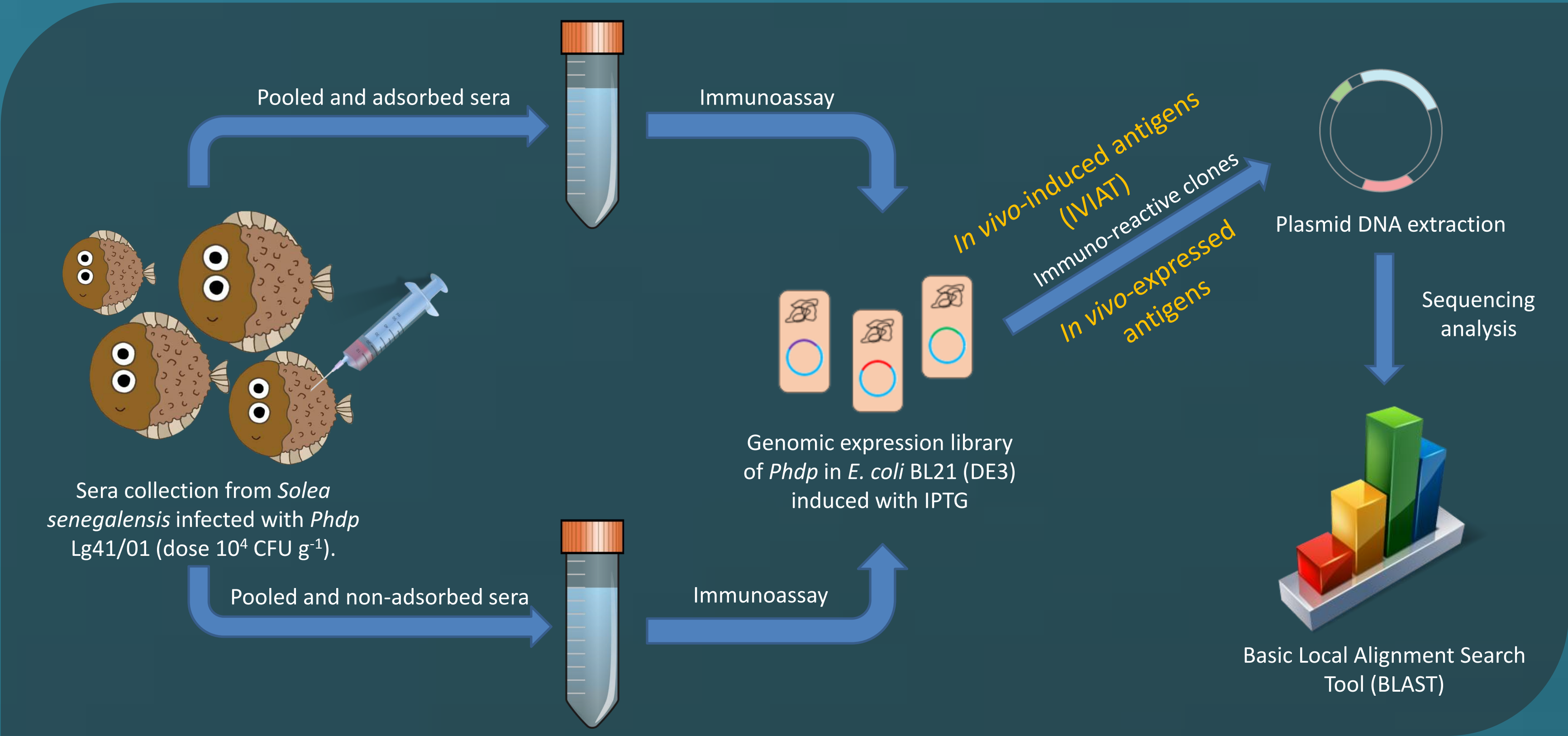
*Photobacterium damsela* subsp. *piscicida* (*Phdp*) is an opportunist pathogen in marine fish responsible for important economic losses. Several virulence factors have been identified in Phd; however, most studies have been carried out *in vitro* and bacterial activities are modulated by their environment. Genes expressed during pathogen infection are important for pathogenicity. In vivo-induced antigen technology (IVIAT) has been used to identify in vivo-induced genes using pooled sera from fish that have experienced photobacteriosis.

## Materials and Methods

Sera were obtained from *S. senegalensis* specimens after sublethal infection with *Phdp* and subsequently pooled and adsorbed against *in vitro* grown *Phdp* Lg41/01 and *Escherichia coli* BL21 (DE3) cells and lysates to remove antibodies against *in vitro* expressed antigens according to Handfield et al. [1]. The efficiency of sera adsorption was evaluated by ELISA, based on the immuno-reactivity with whole and lysed *Phdp* cells, grown *in vitro*, as immobilized antigens. A genomic expression library of *Phdp* Lg41/01 was generated in *E. coli* BL21 (DE3) using pET-30 expression system (Novagen, San Diego, CA, USA). The expression library was probed with adsorbed (for *in vivo*-induced antigens, IVIAT) and non-absorbed sera (for *in vivo*-expressed antigens) using immunoblot technique. Inserted DNA from reactive clones was sequenced (Macrogen Europe, Amsterdam, The Netherlands). Nucleotide sequences were compared against the NCBI protein database using BLASTx.

## Results

Specific antibody titers against *Phdp* were determined in the sera from each surviving *S. senegalensis* specimen. All fish showed significantly higher antibody titers compared to control fish (non-infected fish). After sera adsorption rounds, a progressive reduction in sera immuno-reactivity against *in vitro* grown *Phdp* cells was detected, especially after the first adsorption step. Thus, following adsorption steps substantially resulted in relative enrichment in antibodies recognizing *in vivo* expressed antigens. The library from *Phdp* Lg14/01 constructed in *E. coli* BL21 (DE3) consisted of approximately 6500 recombinants.



Functional category	Identification	Accession	Predicted cellular location
Replication, recombination and repair	DNA gyrase subunit B	WP_044180341.1	Cytoplasm
	Recombination associated protein RdgC	WP_044176475.1	Cytoplasm
Iron acquisition	Non-ribosomal peptide synthetase	AKQ52531	Membrane
Transport and metabolism	Inosine-5'-monophosphate dehydrogenase	EEZ41661.1	Cytoplasm
	Arginine decarboxylase catabolic	AEU10010.1	Cytoplasm
	Glutamine amidotransferases class-II (GATase)	WP_044179637.1	Cytoplasm
	Putative amidotransferase	AEU09986.1	Cytoplasm
	Diguanylate phosphodiesterase	WP_044178285.1	Membrane/ Cytoplasm
Antioxidant activity	Alkyl hydroperoxide reductase	WP_005298372.1	Cytoplasm
	Superoxide dismutase	WP_005298367.1	Periplasm
Cell envelope and wall metabolism	Lytic transglycosylase	WP_044174705.1	Outer cell membrane
	Murein transglycosylase	WP_044178572.1	Outer cell membrane
Translation, ribosomal structure and biogenesis	23S rRNA methyltransferase	WP_044175535.1	Cytoplasm
	Serine hydroxymethyltransferase	WP_005300693.1	Cytoplasm
	Alanyl-tRNA synthetase	WP_044174517.1	Cytoplasm
Cell cycle control, cell division, chromosome partitioning	Chromosome partitioning protein ParA	WP_014386679.1	Cytoplasm
Mobile and extrachromosomal element functions	Putative transposase	AEU10011.1	Cytoplasm
Other	Intramembrane serine protease GlpG	WP_044173910.1	Membrane

*in vivo* compared to *in vitro* conditions might give an insight into the genes relevant to the bacterial virulence [2]. AhpC peroxidase activity has a protective role by reducing hydrogen peroxide, peroxyxynitrite and organic hydroperoxides. Immunization with AhpC conferred protection against *Helicobacter pylori* infection [3]. Impdh catalyzes the conversion of products essential in *de novo* synthesis of guanine nucleotides. Adequate levels of purine nucleotides are critical for cell proliferation, nucleic acid replication, cell signaling and as a biochemical energy source. This gene is an important therapeutic target against bacterial diseases [4]. On the other hand, the non-ribosomal peptide synthetase involved in the synthesis of the siderophore piscibactin is considered a virulence factor in *Phdp*. In the present work, the induction in *S. senegalensis* and its immunogenic character have been determined.

## Conclusion

Different proteins expressed during *Phdp* infection in *S. senegalensis* have been identified. Among them, Impdh, AhpC, Shmt, AlaRS and Irp2 have been identified as *in vivo* induced antigens expressed during *S. senegalensis* infection with *Phdp*. They are likely to play a role in the virulence of *Phdp*. The antigenic character of these proteins makes them potential targets for the development of new vaccines.

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### References

- [1] Handfield, M., Brady, L.J., Progulskie-Fox, A., and Hillman, J.D., 2000. IVIAT: a novel method to identify microbial genes expressed specifically during human infections. Trends in Microbiology, 8: 336-339.
- [2] Li S, Song J, Huang H, Chen W, Li M, Zhao Y, Cong Y, Zhu J, Rao X, Hu X & Hu F. (2013). Identification of *in-vivo* induced genes of *Streptococcus suis* serotype 2 specially expressed in infected human. Microbial Pathogenesis 63: 8-15
- [3] O'Riordan A.A., Morales V.A., Mulligan L., Faheem N., Windle H.J., and Kelleher D.P., 2012. Alkyl hydroperoxide reductase: a candidate *Helicobacter pylori* vaccine. Vaccine, 30:3876-3884.
- [4] Shu, Q., and Nair, V., 2008. Inosine monophosphate dehydrogenase (IMPDH) as a target in drug discovery. Medicinal Research Reviews, 28:219-232.